**RAW SEQUENCE LISTING**
PATENT APPLICATION: **US/10/038,297**

DATE: 12/26/2002
TIME: 15:14:34

Input Set : **A:\10-038,297.asc.txt**
Output Set: **N:\CRF4\12262002\J038297.raw**

```
  1 <110> APPLICANT: Phan, Brigitte Chau
  2       Virtanen, Jorma Antero
  4 <120> TITLE OF INVENTION: Dual Bead Assays Including Covalent Linkages For
  5       Improved Specificity And Related Optical Analysis Discs
  7 <130> FILE REFERENCE: BTI2 00103501(USP2)US
  9 <140> CURRENT APPLICATION NUMBER: 10/038,297
C--> 11 <141> CURRENT FILING DATE: 2002-12-10
W--> 13 <150> PRIOR APPLICATION NUMBER:
W--> 14 <151> PRIOR FILING DATE:
 16 <160> NUMBER OF SEQ ID NOS: 2
 18 <170> SOFTWARE: Microsoft Notepad Windows98
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 40
 22 <212> TYPE: DNA
 23 <213> ORGANISM: Artificial Sequence
W--> 25 <220> FEATURE: No Biological Significance.
W--> 26 Used for experimental purpose only.
W--> 27 <221> NAME/KEY:
 28 <222> LOCATION:
 29 <223> OTHER INFORMATION: Capture Probe
 31 <400> SEQUENCE: 1
 32 ccagtgaatt cgagctcggt acccggggat cctctagagt     40
 34 <210> SEQ ID NO: 2
 35 <211> LENGTH: 40
 36 <212> TYPE: DNA
 37 <213> ORGANISM: Artificial Sequence
W--> 39 <220> FEATURE: No Biological Significance.
W--> 40 Used for experimental purpose only.
W--> 41 <221> NAME/KEY:
 42 <222> LOCATION:
 43 <223> OTHER INFORMATION: Reporter Probe
 45 <400> SEQUENCE: 2
 46 cttagtcttt agatgcaagc ttggcgtaat catggtcata     40
```

**VERIFICATION SUMMARY**                    DATE: 12/26/2002
PATENT APPLICATION:  **US/10/038,297**      TIME: 15:14:35

Input Set : **A:\10-038,297.asc.txt**
Output Set: **N:\CRF4\12262002\J038297.raw**

```
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:13 M:256 W: Invalid Numeric Header Field, <150> PRIOR APPLICATION NUMBER:
L:14 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:25 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:26 M:259 W: Allowed number of lines exceeded, <220> FEATURE:
L:27 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:39 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:40 M:259 W: Allowed number of lines exceeded, <220> FEATURE:
L:41 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
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